



SEQUENCE LISTING

<110> Clarkson, Kathleen A.
Fenel, Fred

<120> Modified Enzymes, Methods to Produce
Modified Enzymes and Uses Thereof

<130> GC812-C1

<140> US 11/404,460

<141> 2006-04-14

<150> US 10/565,954

<151> 2004-09-10

<150> US 60/503,251

<151> 2003-09-15

<160> 51

<170> FastSEQ for Windows Version 4.0

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<211> 223

<212> PRT

<213> Trichoderma reesei

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Arg	Gln	Thr	Ile	Gln	Pro	Gly	Thr	Gly	Tyr	Asn	Asn	Gly	Tyr	Phe	Tyr
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Ser	Tyr	Trp	Asn	Asp	Gly	His	Gly	Gly	Val	Thr	Tyr	Thr	Asn	Gly	Pro
	50					55					60				
Gly	Gly	Gln	Phe	Ser	Val	Asn	Trp	Ser	Asn	Ser	Gly	Asn	Phe	Val	Gly
65						70				75					80
Gly	Lys	Gly	Trp	Gln	Pro	Gly	Thr	Lys	Asn	Lys	Val	Ile	Asn	Phe	Ser
				85					90					95	
Gly	Ser	Tyr	Asn	Pro	Asn	Gly	Asn	Ser	Tyr	Leu	Ser	Val	Tyr	Gly	Trp
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Ser	Arg	Asn	Pro	Leu	Ile	Glu	Tyr	Tyr	Ile	Val	Glu	Asn	Phe	Gly	Thr
		115					120					125			
Tyr	Asn	Pro	Ser	Thr	Gly	Ala	Thr	Lys	Leu	Gly	Glu	Val	Thr	Ser	Asp
	130					135					140				
Gly	Ser	Val	Tyr	Asp	Ile	Tyr	Arg	Thr	Gln	Arg	Val	Asn	Gln	Pro	Ser
145					150					155					160
Ile	Ile	Gly	Thr	Ala	Thr	Phe	Tyr	Gln	Tyr	Trp	Ser	Val	Arg	Arg	Asn
				165					170					175	
His	Arg	Ser	Ser	Gly	Ser	Val	Asn	Thr	Ala	Asn	His	Phe	Asn	Ala	Trp
			180					185					190		
Ala	Gln	Gln	Gly	Leu	Thr	Leu	Gly	Thr	Met	Asp	Tyr	Gln	Ile	Val	Ala
		195					200						205		

Val Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser
 210 215 220

<210> 2
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 <212> DNA
 <213> Trichoderma reesei

<400> 2
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 cccgccgccg aggtcgaaatc cgtggctgtg gagaagcgcc agacgattca gcccggcacg 120
 ggctacaaca acggctactt ctactcgtae tggaacgatg gccacggcgg cgtgacgtac 180
 accaatgggtc ccggcgggca gttctccgtc aactgggtcca actcggggcaa ctttgtcggc 240
 ggcaagggat ggcagcccgg caccaagaac aagtaagact acctactctt accccctttg 300
 accaacacag cacaacacaa tacaacacat gtgactacca atcatggaat cggatctaac 360
 agctgtgttt tcaaaaaaaaaa gggatcatcaa cttctcgggc agtacaacc ccaacggcaa 420
 cagctacctc tccgtgtacg gctgggtccc caacccccctg atcgagtact acatcgtcga 480
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 cggcagcgtc tacgacattt accgcacgca gcgcgtcaac cagccgtcca tcatcggcac 600
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 cagggcgaac cacttcaacg cgtggggtca gcaaggcctg acgctcggga cgatggatta 720
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<210> 3
 <211> 234
 <212> PRT
 <213> Trichoderma reesei

<400> 3
 Met Lys Phe Leu Gln Val Leu Pro Ala Leu Ile Pro Ala Ala Leu Ala
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 Gln Thr Ser Cys Asp Gln Trp Ala Thr Phe Thr Gly Asn Gly Tyr Thr
 20 25 30
 Val Ser Asn Asn Leu Trp Gly Ala Ser Ala Gly Ser Gly Phe Gly Cys
 35 40 45
 Val Thr Ala Val Ser Leu Ser Gly Gly Ala Ser Trp His Ala Asp Trp
 50 55 60
 Gln Trp Ser Gly Gly Gln Asn Asn Val Lys Ser Tyr Gln Asn Ser Gln
 65 70 75 80
 Ile Ala Ile Pro Gln Lys Arg Thr Val Asn Ser Ile Ser Ser Met Pro
 85 90 95
 Thr Thr Ala Ser Trp Ser Tyr Ser Gly Ser Asn Ile Arg Ala Asn Val
 100 105 110
 Ala Tyr Asp Leu Phe Thr Ala Ala Asn Pro Asn His Val Thr Tyr Ser
 115 120 125
 Gly Asp Tyr Glu Leu Met Ile Trp Leu Gly Lys Tyr Gly Asp Ile Gly
 130 135 140
 Pro Ile Gly Ser Ser Gln Gly Thr Val Asn Val Gly Gly Gln Ser Trp
 145 150 155 160
 Thr Leu Tyr Tyr Gly Tyr Asn Gly Ala Met Gln Val Tyr Ser Phe Val
 165 170 175
 Ala Gln Thr Asn Thr Thr Asn Tyr Ser Gly Asp Val Lys Asn Phe Phe
 180 185 190
 Asn Tyr Leu Arg Asp Asn Lys Gly Tyr Asn Ala Ala Gly Gln Tyr Val
 195 200 205
 Leu Ser Tyr Gln Phe Gly Thr Glu Pro Phe Thr Gly Ser Gly Thr Leu

210
Asn Val Ala Ser Trp Thr Ala Ser Ile Asn
225 230

220

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<212> DNA
<213> Trichoderma reesei

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tcagccggct ctggatttgg ctgctgacg gcggtatcgc tcagcggcgg ggctcctgg 180
cacgcagact ggcagtggc cggcggccag aacaacgtca agtcgtacca gaactctcag 240
attgccattc cccagaagag gaccgtcaac agcatcagca gcatgccac cactgccagc 300
tggagctaca gcgggagcaa catccgcgct aatgttgcgt atgacttggt caccgcagcc 360
aaccggaatc atgtcacgta ctcgggagac tacgaactca tgatctggta agccataaga 420
agtgaccctc cttgatagtt tcgactaaca acatgtcttg aggcttggca aatacggcga 480
tattgggccc attgggtcct cacagggaac agtcaacgtc ggtggccaga gctggacgct 540
ctactatggc tacaacggag ccatgcaagt ctattccttt gtggcccaga ccaacactac 600
caactacagc ggagatgtca agaacttctt caattatctc cgagacaata aaggatacaa 660
cgctgcaggc caatatgttc ttagtaagtc accctcactg tgactgggct gagtttggtg 720
caacgtttgc taacaaaacc ttcgtatagg ctaccaattt ggtaccgagc ccttcacggg 780
cagtgggaact ctgaacgtcg catcctggac cgcattctat aactaa 826

<210> 5
<211> 222
<212> PRT
<213> Trichoderma reesei

<400> 5
Met Val Ser Phe Thr Ser Leu Leu Ala Ala Ser Pro Pro Ser Arg Ala
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Ser Cys Arg Pro Ala Ala Glu Val Glu Ser Val Ala Val Glu Lys Arg
20 25 30
Gln Thr Ile Gln Pro Gly Thr Gly Tyr Asn Asn Gly Tyr Phe Tyr Ser
35 40 45
Tyr Trp Asn Asp Gly His Gly Gly Val Thr Tyr Thr Asn Gly Pro Gly
50 55 60
Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly
65 70 75 80
Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly
85 90 95
Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser
100 105 110
Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr
115 120 125
Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly
130 135 140
Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile
145 150 155 160
Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His
165 170 175
Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala
180 185 190
Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val
195 200 205

Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser
 210 215 220

<210> 6
 <211> 227
 <212> PRT
 <213> Humicola insolens

<400> 6
 Met Val Ser Leu Lys Ser Val Leu Ala Ala Ala Thr Ala Val Ser Ser
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 Ala Ile Ala Ala Pro Phe Asp Phe Val Pro Arg Asp Asn Ser Thr Ala
 20 25 30
 Leu Gln Ala Arg Gln Val Thr Pro Asn Ala Glu Gly Trp His Asn Gly
 35 40 45
 Tyr Phe Tyr Ser Trp Trp Ser Asp Gly Gly Gly Gln Val Gln Tyr Thr
 50 55 60
 Asn Leu Glu Gly Ser Arg Tyr Gln Val Arg Trp Arg Asn Thr Gly Asn
 65 70 75 80
 Phe Val Gly Gly Lys Gly Trp Asn Pro Gly Thr Gly Arg Thr Ile Asn
 85 90 95
 Tyr Gly Gly Tyr Phe Asn Pro Gln Gly Asn Gly Tyr Leu Ala Val Tyr
 100 105 110
 Gly Trp Thr Arg Asn Pro Leu Val Glu Tyr Tyr Val Ile Glu Ser Tyr
 115 120 125
 Gly Thr Tyr Asn Pro Gly Ser Gln Ala Gln Tyr Lys Gly Thr Phe Tyr
 130 135 140
 Thr Asp Gly Asp Gln Tyr Asp Ile Phe Val Ser Thr Arg Tyr Asn Gln
 145 150 155 160
 Pro Ser Ile Asp Gly Thr Arg Thr Phe Gln Tyr Trp Ser Ile Arg
 165 170 175
 Lys Asn Lys Arg Val Gly Gly Ser Val Asn Met Gln Asn His Phe Asn
 180 185 190
 Ala Trp Gln Gln His Gly Met Pro Leu Gly Gln His Tyr Tyr Gln Val
 195 200 205
 Val Ala Thr Glu Gly Tyr Gln Ser Ser Gly Glu Ser Asp Ile Tyr Val
 210 215 220
 Gln Thr His
 225

<210> 7
 <211> 210
 <212> PRT
 <213> Bacillus stearothermophilus

<400> 7
 Met Lys Leu Lys Lys Lys Met Leu Thr Leu Leu Leu Thr Ala Ser Met
 1 5 10 15
 Ser Phe Gly Leu Phe Gly Ala Thr Ser Ser Ala Ala Thr Asp Tyr Trp
 20 25 30
 Gln Tyr Trp Thr Asp Gly Gly Gly Met Val Asn Ala Val Asn Gly Pro
 35 40 45
 Gly Gly Asn Tyr Ser Val Thr Trp Gln Asn Thr Gly Asn Phe Val Val
 50 55 60
 Gly Lys Gly Trp Thr Val Gly Ser Pro Asn Arg Val Ile Asn Tyr Asn
 65 70 75 80
 Ala Gly Ile Trp Glu Pro Ser Gly Asn Gly Tyr Leu Thr Leu Tyr Gly

				85					90					95			
Trp	Thr	Arg	Asn	Ala	Leu	Ile	Glu	Tyr	Tyr	Val	Val	Asp	Ser	Trp	Gly		
			100					105					110				
Thr	Tyr	Arg	Pro	Thr	Gly	Asn	Tyr	Lys	Gly	Thr	Val	Asn	Ser	Asp	Gly		
		115					120					125					
Gly	Thr	Tyr	Asp	Ile	Tyr	Thr	Thr	Met	Arg	Tyr	Asn	Ala	Pro	Ser	Ile		
	130					135					140						
Asp	Gly	Thr	Gln	Thr	Phe	Gln	Gln	Phe	Trp	Ser	Val	Arg	Gln	Ser	Lys		
145					150					155					160		
Arg	Pro	Thr	Gly	Ser	Asn	Val	Ser	Ile	Thr	Phe	Ser	Asn	His	Val	Asn		
			165					170					175				
Ala	Trp	Arg	Ser	Lys	Gly	Met	Asn	Leu	Gly	Ser	Ser	Trp	Ala	Tyr	Gln		
		180						185					190				
Val	Leu	Ala	Thr	Glu	Gly	Tyr	Gln	Ser	Ser	Gly	Arg	Ser	Asn	Val	Thr		
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Val	Trp																
	210																

<210> 8
 <211> 229
 <212> PRT
 <213> Trichoderma reesei

<400> 8																	
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Thr	Leu	Ala	Met	Pro	Thr	Gly	Leu	Glu	Pro	Glu	Ser	Ser	Val	Asn	Val		
		20						25					30				
Thr	Glu	Arg	Gly	Met	Tyr	Asp	Phe	Val	Leu	Gly	Ala	His	Asn	Asp	His		
		35				40						45					
Arg	Arg	Arg	Ala	Ser	Ile	Asn	Tyr	Asp	Gln	Asn	Tyr	Gln	Thr	Gly	Gly		
	50				55					60							
Gln	Val	Ser	Tyr	Ser	Pro	Ser	Asn	Thr	Gly	Phe	Ser	Val	Asn	Trp	Asn		
65				70						75					80		
Thr	Gln	Asp	Asp	Phe	Val	Val	Gly	Val	Gly	Trp	Thr	Thr	Gly	Ser	Ser		
			85					90					95				
Ala	Pro	Ile	Asn	Phe	Gly	Gly	Ser	Phe	Ser	Val	Asn	Ser	Gly	Thr	Gly		
		100						105					110				
Leu	Leu	Ser	Val	Tyr	Gly	Trp	Ser	Thr	Asn	Pro	Leu	Val	Glu	Tyr	Tyr		
		115					120					125					
Ile	Met	Glu	Asp	Asn	His	Asn	Tyr	Pro	Ala	Gln	Gly	Thr	Val	Lys	Gly		
	130			135							140						
Thr	Val	Thr	Ser	Asp	Gly	Ala	Thr	Tyr	Thr	Ile	Trp	Glu	Asn	Thr	Arg		
145				150				155						160			
Val	Asn	Glu	Pro	Ser	Ile	Gln	Gly	Thr	Ala	Thr	Phe	Asn	Gln	Tyr	Ile		
			165					170					175				
Ser	Val	Arg	Asn	Ser	Pro	Arg	Thr	Ser	Gly	Thr	Val	Thr	Val	Gln	Asn		
		180						185					190				
His	Phe	Asn	Ala	Trp	Ala	Ser	Leu	Gly	Leu	His	Leu	Gly	Gln	Met	Asn		
	195					200						205					
Tyr	Gln	Val	Val	Ala	Val	Glu	Gly	Trp	Gly	Gly	Ser	Gly	Ser	Ala	Ser		
	210					215					220						
Gln	Ser	Val	Ser	Asn													
225																	

<210> 9
 <211> 211

<212> PRT

<213> *Aspergillus awamori*

<400> 9

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Met Lys Val Thr Ala Ala Phe Ala Gly Leu Leu Val Thr Ala Phe Ala
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Ala Pro Val Pro Glu Pro Val Leu Val Ser Arg Ser Ala Gly Ile Asn
          20          25          30
Tyr Val Gln Asn Tyr Asn Gly Asn Leu Gly Asp Phe Thr Tyr Asp Glu
          35          40          45
Ser Ala Gly Thr Phe Ser Met Tyr Trp Glu Asp Gly Val Ser Ser Asp
          50          55          60
Phe Val Val Gly Leu Gly Trp Thr Thr Gly Ser Ser Asn Ala Ile Thr
65          70          75          80
Tyr Ser Ala Glu Tyr Ser Ala Ser Gly Ser Ser Ser Tyr Leu Ala Val
          85          90          95
Tyr Gly Trp Val Asn Tyr Pro Gln Ala Glu Tyr Tyr Ile Val Glu Asp
          100          105          110
Tyr Gly Asp Tyr Asn Pro Cys Ser Ser Ala Thr Ser Leu Gly Thr Val
          115          120          125
Tyr Ser Asp Gly Ser Thr Tyr Gln Val Cys Thr Asp Thr Arg Thr Asn
          130          135          140
Glu Pro Ser Ile Thr Gly Thr Ser Thr Phe Thr Gln Tyr Phe Ser Val
145          150          155          160
Arg Glu Ser Thr Arg Thr Ser Gly Thr Val Thr Val Ala Asn His Phe
          165          170          175
Asn Phe Trp Ala Gln His Gly Phe Gly Asn Ser Asp Phe Asn Tyr Gln
          180          185          190
Val Met Ala Val Glu Ala Trp Ser Gly Ala Gly Ser Ala Ser Val Thr
          195          200          205
Ile Ser Ser
          210
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<210> 10

<211> 330

<212> PRT

<213> *Bacillus stearothermophilus*

<400> 10

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Arg Ile Gly Ala Ala Val Asn Pro Val Thr Leu Glu Ala Gln Gln Ser
          20          25          30
Leu Leu Ile Arg His Val Asn Ser Leu Thr Ala Glu Asn His Met Lys
          35          40          45
Phe Glu His Leu Gln Pro Glu Glu Gly Arg Phe Thr Phe Asp Ile Ala
          50          55          60
Ile Lys Ser Ser Thr Ser Pro Phe Ser Ser His Gly Val Arg Gly His
65          70          75          80
Thr Leu Val Trp His Asn Gln Thr Pro Ser Trp Val Phe Gln Asp Ser
          85          90          95
Gln Gly His Phe Val Gly Arg Asp Val Leu Leu Glu Arg Met Lys Ser
          100          105          110
His Ile Ser Thr Val Val Gln Arg Tyr Lys Gly Lys Val Tyr Cys Trp
          115          120          125
Asp Val Ile Asn Glu Ala Val Ala Asp Glu Gly Ser Glu Trp Leu Arg
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130		135		140
Ser Ser Thr Trp Arg Gln Ile Ile Gly Asp Asp Phe Ile Gln Gln Ala				
145		150		155
Phe Leu Tyr Ala His Glu Ala Asp Pro Glu Ala Leu Leu Phe Tyr Asn				160
	165		170	175
Asp Tyr Asn Glu Cys Phe Pro Glu Lys Arg Glu Lys Ile Tyr Thr Leu				
	180		185	190
Val Lys Ser Leu Arg Asp Lys Gly Ile Pro Ile His Gly Ile Gly Met				
	195		200	205
Gln Ala His Trp Ser Leu Asn Arg Pro Thr Leu Asp Glu Ile Arg Ala				
	210		215	220
Ala Ile Glu Arg Tyr Ala Ser Leu Gly Val Ile Leu His Ile Thr Glu				
225		230		235
Leu Asp Ile Ser Met Phe Glu Phe Asp Asp His Arg Lys Asp Leu Ala				240
	245		250	255
Ala Pro Thr Asn Glu Met Val Glu Arg Gln Ala Glu Arg Tyr Glu Gln				
	260		265	270
Ile Phe Ser Leu Phe Lys Glu Tyr Arg Asp Val Ile Gln Asn Val Thr				
	275		280	285
Phe Trp Gly Ile Ala Asp Asp His Thr Trp Leu Asp His Phe Pro Val				
	290		295	300
Gln Gly Arg Lys Asn Trp Pro Leu Leu Phe Asp Glu Gln His Asn Pro				
305		310		315
Lys Pro Ala Phe Trp Arg Val Val Asn Ile				320
	325		330	

<210> 11

<211> 190

<212> PRT

<213> *Trichoderma reesei*

<400> 11

Gln Thr Ile Gln Pro Gly Thr Gly Tyr Asn Asn Gly Tyr Phe Tyr Ser				
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Tyr Trp Asn Asp Gly His Gly Gly Val Thr Tyr Thr Asn Gly Pro Gly				
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Gly Gln Phe Ser Val Asn Trp Ser Asn Ser Gly Asn Phe Val Gly Gly				
	35		40	45
Lys Gly Trp Gln Pro Gly Thr Lys Asn Lys Val Ile Asn Phe Ser Gly				
	50		55	60
Ser Tyr Asn Pro Asn Gly Asn Ser Tyr Leu Ser Val Tyr Gly Trp Ser				
65		70		75
Arg Asn Pro Leu Ile Glu Tyr Tyr Ile Val Glu Asn Phe Gly Thr Tyr				
	85		90	95
Asn Pro Ser Thr Gly Ala Thr Lys Leu Gly Glu Val Thr Ser Asp Gly				
	100		105	110
Ser Val Tyr Asp Ile Tyr Arg Thr Gln Arg Val Asn Gln Pro Ser Ile				
	115		120	125
Ile Gly Thr Ala Thr Phe Tyr Gln Tyr Trp Ser Val Arg Arg Asn His				
	130		135	140
Arg Ser Ser Gly Ser Val Asn Thr Ala Asn His Phe Asn Ala Trp Ala				
145		150		155
Gln Gln Gly Leu Thr Leu Gly Thr Met Asp Tyr Gln Ile Val Ala Val				
	165		170	175
Glu Gly Tyr Phe Ser Ser Gly Ser Ala Ser Ile Thr Val Ser				
	180		185	190

<210> 12
 <211> 237
 <212> PRT
 <213> *Aspergillus awamori*

<400> 12
 Met Lys Ala Phe His Leu Leu Ala Ala Leu Ser Gly Ala Ala Val Ala
 1 5 10 15
 Gln Gln Ala Gln Leu Cys Asp Gln Tyr Ala Thr Tyr Thr Gly Gly Val
 20 25 30
 Tyr Thr Ile Asn Asn Asn Leu Trp Gly Lys Asp Ala Gly Ser Gly Ser
 35 40 45
 Gln Cys Thr Thr Val Asn Ser Ala Ser Ser Ala Gly Thr Ser Trp Ser
 50 55 60
 Thr Lys Trp Asn Trp Ser Gly Gly Glu Asn Ser Val Lys Ser Tyr Ala
 65 70 75 80
 Asn Ser Gly Leu Ser Phe Asn Lys Lys Leu Val Ser Gln Ile Ser His
 85 90 95
 Ile Pro Thr Ala Ala Arg Trp Ser Tyr Asp Asn Thr Cys Ile Arg Arg
 100 105 110
 Gly Arg Ala Tyr Asp Leu Phe Thr Ala Ala Asp Ile Asn His Val Thr
 115 120 125
 Trp Ser Gly Asp Tyr Glu Leu Met Ile Trp Leu Ala Arg Tyr Gly Gly
 130 135 140
 Val Gln Pro Leu Gly Ser Gln Ile Ala Thr Ala Thr Val Glu Gly Gln
 145 150 155 160
 Thr Trp Glu Leu Trp Tyr Gly Val Asn Gly Ala Gln Lys Thr Tyr Ser
 165 170 175
 Phe Val Ala Ala Asn Pro Ile Thr Ser Phe Gln Gly Asp Ile Asn Asp
 180 185 190
 Phe Phe Lys Tyr Leu Thr Gln Asn His Gly Phe Pro Ala Ser Ser Gln
 195 200 205
 Tyr Leu Ile Thr Leu Gln Phe Gly Thr Glu Pro Phe Thr Gly Gly Pro
 210 215 220
 Ala Thr Leu Asn Val Ala Asp Trp Ser Ala Ser Val Gln
 225 230 235

<210> 13
 <211> 233
 <212> PRT
 <213> *Trichoderma viride*

<400> 13
 Met Lys Phe Leu Gln Ile Ala Pro Thr Leu Leu Pro Val Ala Leu Ala
 1 5 10 15
 Gln Ser Ser Cys Ser Gln Tyr Ala Thr Phe Ser Gly Gly Asn Tyr Ala
 20 25 30
 Leu Ser Asn Asn Leu Trp Gly Gln Ser Ala Gly Ser Gly Ser Gly Cys
 35 40 45
 Ile Thr Asp Val Ser Leu Gly Gly Ser Ala Val Trp Ser Thr Thr Trp
 50 55 60
 Asp Trp Ser Gly Gly Gln Ser Asn Val Lys Gly Tyr Pro Asn Ile Ala
 65 70 75 80
 Leu Asn Ile Pro Asn Lys Arg Leu Val Ser Ser Ile Ser Ser Met Pro
 85 90 95
 Thr Thr Ala Gln Trp Ser Tyr Ser Gly Ser Ser Ile Arg Ala Asp Val

			100						105				110			
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ser	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser	
		115						120					125			
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Tyr	Gly	Asp	Ile	Gln	
	130					135					140					
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Thr	Ser	Trp	
145					150					155					160	
Asn	Leu	Trp	Tyr	Gly	Pro	Asn	Gly	Ser	Met	Gln	Val	Tyr	Ser	Phe	Val	
				165					170					175		
Ala	Pro	Gly	Asn	Leu	Thr	Asn	Trp	Ser	Gly	Asp	Val	Lys	Asn	Phe	Tyr	
		180						185					190			
Thr	Tyr	Leu	Gln	Asn	Asn	Lys	Gly	Tyr	Pro	Ala	Ser	Ser	Gln	Tyr	Val	
	195						200					205				
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Ala	Phe	Thr	Gly	Ser	Gly	Thr	Leu	
	210					215					220					
Asn	Asn	Thr	Trp	Thr	Ala	Ser	Ile	Asn								
225					230											

<210> 14
 <211> 234
 <212> PRT
 <213> Hypocrea koningii

Met	Lys	Leu	Ile	His	Val	Leu	Pro	Ala	Leu	Ile	Pro	Ala	Ala	Leu	Ala	
1				5					10					15		
Gln	Thr	Ser	Cys	Asp	Gln	Tyr	Ala	Val	Phe	Thr	Gly	Ser	Asp	Tyr	Thr	
			20					25					30			
Val	Ser	Asn	Asn	Leu	Trp	Gly	Gln	Ser	Ala	Gly	Ser	Gly	Phe	Gly	Cys	
	35					40						45				
Val	Thr	Ala	Glu	Ser	Leu	Ser	Gly	Ser	Ala	Ser	Trp	His	Ala	Asp	Trp	
	50					55					60					
Gln	Trp	Ser	Gly	Gly	Gln	Asn	Asn	Val	Lys	Ser	Tyr	Gln	Asn	Ser	Gln	
65					70					75					80	
Ile	Pro	Ile	Pro	Gln	Lys	Arg	Thr	Val	Asn	Ser	Ile	Ser	Ser	Met	Pro	
				85				90						95		
Thr	Thr	Ala	Ser	Trp	Ser	Tyr	Thr	Gly	Ser	Asp	Ile	Arg	Ala	Asn	Val	
		100						105					110			
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser	
	115					120						125				
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Arg	Tyr	Gly	Asp	Ile	Gly	
	130					135					140					
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Gln	Ser	Trp	
145					150					155					160	
Thr	Leu	Tyr	Tyr	Gly	Tyr	Asn	Gly	Ala	Met	Gln	Val	Tyr	Ser	Phe	Val	
				165				170						175		
Ala	Gln	Thr	Asn	Thr	Thr	Ser	Tyr	Ser	Gly	Asp	Val	Lys	Asn	Phe	Phe	
		180						185					190			
Asn	Tyr	Leu	Arg	Asp	Asn	Lys	Gly	Tyr	Asn	Ala	Ala	Gly	Gln	Tyr	Val	
	195						200					205				
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Gly	Thr	Leu	
	210					215					220					
Asn	Val	Ala	Ser	Trp	Thr	Ala	Ser	Ile	Asn							
225					230											

<210> 15
 <211> 234

<212> PRT

<213> *Hypocrea schweinitzii*

<400> 15

Met	Lys	Phe	Leu	Gln	Val	Leu	Pro	Ala	Ile	Leu	Pro	Ala	Ala	Leu	Ala
1				5					10					15	
Gln	Thr	Ser	Cys	Asp	Gln	Tyr	Ala	Thr	Phe	Ser	Gly	Asn	Gly	Tyr	Ile
			20					25					30		
Val	Ser	Asn	Asn	Leu	Trp	Gly	Ala	Ser	Ala	Gly	Ser	Gly	Phe	Gly	Cys
		35					40					45			
Val	Thr	Ser	Val	Ser	Leu	Asn	Gly	Ala	Ala	Ser	Trp	His	Ala	Asp	Trp
	50					55					60				
Gln	Trp	Ser	Gly	Gly	Gln	Asn	Asn	Val	Lys	Ser	Tyr	Gln	Asn	Val	Gln
65					70					75				80	
Ile	Asn	Ile	Pro	Gln	Lys	Arg	Thr	Val	Asn	Ser	Ile	Gly	Ser	Met	Pro
				85					90					95	
Thr	Thr	Ala	Ser	Trp	Ser	Tyr	Ser	Gly	Ser	Asp	Ile	Arg	Ala	Asn	Val
			100					105					110		
Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Val	Thr	Tyr	Ser
		115					120					125			
Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Tyr	Gly	Asp	Ile	Gly
	130					135					140				
Pro	Ile	Gly	Ser	Ser	Gln	Gly	Thr	Val	Asn	Val	Gly	Gly	Gln	Thr	Trp
145					150					155				160	
Thr	Leu	Tyr	Tyr	Gly	Tyr	Asn	Gly	Ala	Met	Gln	Val	Tyr	Ser	Phe	Val
				165				170						175	
Ala	Gln	Ser	Asn	Thr	Thr	Ser	Tyr	Ser	Gly	Asp	Val	Lys	Asn	Phe	Phe
			180					185					190		
Asn	Tyr	Leu	Arg	Asp	Asn	Lys	Gly	Tyr	Asn	Ala	Gly	Gly	Gln	Tyr	Val
	195						200					205			
Leu	Ser	Tyr	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Gly	Thr	Leu
	210					215					220				
Asn	Val	Ala	Ser	Trp	Thr	Ala	Ser	Ile	Asn						
225					230										

<210> 16

<211> 237

<212> PRT

<213> *Stachybotrys echinata*

<400> 16

Met	Lys	Val	Ala	Ala	Leu	Leu	Val	Ala	Leu	Ser	Pro	Leu	Ala	Phe	Ala
1				5					10					15	
Gln	Ser	Leu	Cys	Asp	Gln	Tyr	Ser	Tyr	Tyr	Ser	Ser	Asn	Gly	Tyr	Glu
			20					25					30		
Phe	Asn	Asn	Asn	Met	Trp	Gly	Arg	Asn	Ser	Gly	Gln	Gly	Asn	Gln	Cys
		35					40					45			
Thr	Tyr	Val	Asp	Tyr	Ser	Ser	Pro	Asn	Gly	Val	Gly	Trp	Arg	Val	Asn
	50					55					60				
Trp	Asn	Trp	Ser	Gly	Gly	Asp	Asn	Asn	Val	Lys	Ser	Tyr	Pro	Tyr	Ser
65					70					75				80	
Gly	Arg	Gln	Leu	Pro	Thr	Lys	Arg	Ile	Val	Ser	Trp	Ile	Gly	Ser	Leu
				85					90					95	
Pro	Thr	Thr	Val	Ser	Trp	Asn	Tyr	Gln	Gly	Asn	Asn	Leu	Arg	Ala	Asn
			100					105					110		
Val	Ala	Tyr	Asp	Leu	Phe	Thr	Ala	Ala	Asn	Pro	Asn	His	Pro	Asn	Ser
		115					120					125			

Ser Gly Asp Tyr Glu Leu Met Ile Trp Leu Gly Arg Leu Gly Asn Val
 130 135 140
 Tyr Pro Ile Gly Asn Gln Val Ala Thr Val Asn Ile Ala Gly Gln Gln
 145 150 155 160
 Trp Asn Leu Tyr Tyr Gly Tyr Asn Gly Ala Met Gln Val Tyr Ser Phe
 165 170 175
 Val Ser Pro Asn Gln Leu Asn Tyr Phe Ser Gly Asn Val Lys Asp Phe
 180 185 190
 Phe Thr Tyr Leu Gln Tyr Asn Arg Ala Tyr Pro Ala Asp Ser Gln Tyr
 195 200 205
 Leu Ile Thr Tyr Gln Phe Gly Thr Glu Pro Phe Thr Gly Gln Asn Ala
 210 215 220
 Val Phe Thr Val Ser Asn Trp Ser Ala Gln Gln Asn Asn
 225 230 235

<210> 17

<211> 238

<212> PRT

<213> *Fusarium equiseti*

<400> 17

Met Lys Ser Thr Leu Leu Leu Ala Gly Ala Phe Ala Pro Leu Ala Phe
 1 5 10 15
 Ala Lys Asp Leu Cys Glu Gln Tyr Gly Tyr Leu Ser Ser Asp Gly Tyr
 20 25 30
 Ser Leu Asn Asn Asn Val Trp Gly Lys Asp Ser Gly Thr Gly Asp Gln
 35 40 45
 Cys Thr His Val Asn Trp Asn Asn Ala Asn Gly Ala Gly Trp Asp Val
 50 55 60
 Glu Trp Asn Trp Ser Gly Gly Lys Asp Asn Val Lys Ser Tyr Pro Asn
 65 70 75 80

Ser Ala Leu Leu Ile Gly Glu Asp Lys Lys Thr Ile Ser Ser Ile Thr
 85 90 95
 Asn Met Gln Ser Thr Ala Glu Trp Lys Tyr Ser Gly Asp Asn Leu Arg
 100 105 110
 Ala Asp Val Ala Tyr Asp Leu Phe Thr Ala Ala Asp Pro Asn His Glu
 115 120 125
 Thr Ser Ser Gly Glu Tyr Glu Leu Met Val Trp Leu Ala Arg Ile Gly
 130 135 140
 Gly Val Gln Pro Ile Gly Ser Leu Gln Thr Ser Val Thr Ile Glu Gly
 145 150 155 160
 His Thr Trp Glu Leu Trp Val Gly Met Asn Gly Ser Met Lys Val Phe
 165 170 175
 Ser Phe Val Ala Pro Thr Pro Val Asn Asn Phe Asn Ala Asp Ile Lys
 180 185 190
 Gln Phe Trp Asp Tyr Leu Thr Lys Ser Gln Asn Phe Pro Ala Asp Asn
 195 200 205
 Gln Tyr Leu Leu Thr Phe Gln Phe Gly Thr Glu Pro Phe Thr Gly Asp
 210 215 220
 Asn Ala Lys Phe Thr Val Thr Asn Phe Asn Ala His Leu Lys
 225 230 235

<210> 18

<211> 237

<212> PRT

<213> *Bionectria ochroleuca*

<400> 18

Met	Lys	Thr	Gly	Ile	Ala	Tyr	Leu	Ala	Ala	Val	Leu	Pro	Leu	Ala	Met
1				5				10						15	
Ala	Glu	Ser	Leu	Cys	Asp	Gln	Tyr	Ala	Tyr	Leu	Ser	Arg	Asp	Gly	Tyr
			20					25					30		
Asn	Phe	Asn	Asn	Asn	Glu	Trp	Gly	Ala	Ala	Thr	Gly	Thr	Gly	Asp	Gln
		35					40					45			
Cys	Thr	Tyr	Val	Asp	Ser	Thr	Ser	Ser	Gly	Gly	Val	Ser	Trp	His	Ser
	50					55					60				
Asp	Trp	Thr	Asn	Ser	Gly	Ser	Glu	Ser	Glu	Ile	Lys	Ser	Tyr	Pro	Tyr
65					70					75					80
Ser	Gly	Leu	Asp	Leu	Pro	Glu	Lys	Lys	Ile	Val	Thr	Ser	Ile	Gly	Ser
				85					90					95	
Ile	Ser	Thr	Gly	Ala	Glu	Trp	Ser	Tyr	Ser	Gly	Ser	Asn	Ile	Arg	Ala
			100					105					110		
Asp	Val	Ala	Tyr	Asp	Ile	Phe	Thr	Ala	Ala	Asp	Pro	Asn	His	Ala	Thr
		115					120					125			
Ser	Ser	Gly	Asp	Tyr	Glu	Val	Met	Ile	Trp	Leu	Ala	Asn	Leu	Gly	Gly
		130				135					140				
Leu	Thr	Pro	Ile	Gly	Ser	Pro	Ile	Gly	Thr	Val	Lys	Ala	Ala	Gly	Arg
145				150						155					160
Asp	Trp	Glu	Leu	Trp	Asp	Gly	Tyr	Asn	Gly	Ala	Met	Arg	Val	Tyr	Ser
				165					170					175	
Phe	Val	Ala	Pro	Ser	Gln	Leu	Asn	Ser	Phe	Asp	Gly	Glu	Ile	Met	Asp
		180					185						190		
Phe	Phe	Tyr	Val	Val	Lys	Asp	Met	Arg	Gly	Phe	Pro	Ala	Asp	Ser	Gln
		195				200						205			
His	Leu	Leu	Thr	Val	Gln	Phe	Gly	Thr	Glu	Pro	Ile	Ser	Gly	Ser	Gly
	210				215						220				
Ala	Lys	Phe	Ser	Val	Ser	His	Trp	Ser	Ala	Lys	Leu	Gly			
225					230					235					

<210> 19

<211> 236

<212> PRT

<213> Bionectria ochroleuca

<400> 19

Met	Lys	Phe	Gln	Leu	Leu	Ser	Leu	Thr	Ala	Phe	Ala	Pro	Leu	Ser	Leu
1				5					10					15	
Ala	Ala	Leu	Cys	Gly	Gln	Tyr	Gln	Ser	Gln	Ser	Gln	Gly	Gly	Tyr	Ile
			20					25					30		
Phe	Asn	Asn	Asn	Lys	Trp	Gly	Gln	Gly	Ser	Gly	Ser	Gly	Ser	Gln	Cys
		35					40					45			
Leu	Thr	Ile	Asp	Lys	Thr	Trp	Asp	Ser	Asn	Val	Ala	Phe	His	Ala	Asp
	50					55					60				
Trp	Ser	Trp	Ser	Gly	Gly	Thr	Asn	Asn	Val	Lys	Ser	Tyr	Pro	Asn	Ala
65					70					75					80
Gly	Leu	Glu	Phe	Ser	Arg	Gly	Lys	Lys	Val	Ser	Ser	Ile	Gly	Thr	Ile
				85					90					95	
Asn	Gly	Gly	Ala	Asp	Trp	Asp	Tyr	Ser	Gly	Ser	Asn	Ile	Arg	Ala	Asn
			100					105					110		
Val	Ala	Tyr	Asp	Ile	Phe	Thr	Ser	Ala	Asp	Pro	Asn	His	Val	Thr	Ser
		115					120					125			
Ser	Gly	Asp	Tyr	Glu	Leu	Met	Ile	Trp	Leu	Gly	Lys	Leu	Gly	Asp	Ile
	130					135					140				

Tyr	Pro	Ile	Gly	Asn	Ser	Ile	Gly	Arg	Val	Lys	Ala	Ala	Asn	Arg	Glu
145					150					155					160
Trp	Asp	Leu	His	Val	Gly	Tyr	Asn	Gly	Ala	Met	Lys	Val	Phe	Ser	Phe
				165					170					175	
Val	Ala	Pro	Ser	Pro	Val	Thr	Arg	Phe	Asp	Gly	Asn	Ile	Met	Asp	Phe
			180					185					190		
Phe	Tyr	Val	Met	Arg	Asp	Met	Gln	Gly	Tyr	Pro	Met	Asp	Lys	Gln	Tyr
		195					200					205			
Leu	Leu	Thr	Leu	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser	Asn	Ala
		210				215					220				
Lys	Phe	Ser	Cys	Trp	Tyr	Phe	Gly	Ala	Lys	Ile	Lys				
225					230					235					

<210> 20

<211> 240

<212> PRT

<213> Bionectria ochroleuca

<400> 20

Met	Lys	Ala	Asn	Ile	Val	Ile	Leu	Ser	Leu	Phe	Ala	Pro	Leu	Ala	Ala
1				5					10					15	
Val	Ala	Gln	Thr	Leu	Cys	Gly	Gln	Tyr	Ser	Ser	Asn	Thr	Gln	Gly	Gly
			20					25					30		
Tyr	Ile	Phe	Asn	Asn	Asn	Met	Trp	Gly	Met	Gly	Ser	Gly	Ser	Gly	Ser
		35					40					45			
Gln	Cys	Thr	Tyr	Val	Asp	Lys	Val	Trp	Ala	Glu	Gly	Val	Ala	Trp	His
	50					55					60				
Thr	Asp	Trp	Ser	Trp	Ser	Gly	Gly	Asp	Asn	Asn	Val	Lys	Ser	Tyr	Pro
65					70				75					80	
Tyr	Ser	Gly	Arg	Glu	Leu	Gly	Thr	Lys	Arg	Ile	Val	Ser	Ser	Ile	Lys
				85				90						95	
Ser	Ile	Ser	Ser	Gly	Ala	Asp	Trp	Asp	Tyr	Thr	Gly	Ser	Asn	Leu	Arg
			100					105					110		
Ala	Asn	Ala	Ala	Tyr	Asp	Ile	Phe	Thr	Ser	Ala	Asn	Pro	Asn	His	Ala
		115					120					125			
Thr	Ser	Ser	Gly	Asp	Tyr	Glu	Val	Met	Ile	Trp	Leu	Gly	Arg	Tyr	Gly
	130					135					140				
Gly	Val	Tyr	Pro	Ile	Gly	Asn	Ser	Ile	Gly	Thr	Val	Arg	Ala	Ala	Gly
145					150					155					160
Arg	Asp	Trp	Ala	Leu	His	Ile	Gly	Tyr	Asn	Gly	Ala	Met	Lys	Val	Phe
				165					170					175	
Ser	Phe	Val	Ala	Ala	Asn	Pro	Val	Thr	Arg	Phe	Asp	Gly	Glu	Ile	Met
			180					185					190		
Asp	Phe	Phe	Tyr	Leu	Leu	Arg	Asp	Met	Gln	Gly	Tyr	Pro	Met	Thr	Ser
		195					200					205			
Gln	Tyr	Leu	Leu	Thr	Leu	Gln	Phe	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Ser
	210					215					220				
Gly	Ala	Lys	Phe	Asn	Cys	Trp	Tyr	Phe	Gly	Ala	Thr	Leu	Ser	Tyr	Trp
225					230					235					240

<210> 21

<211> 254

<212> PRT

<213> Humicola grisea

<400> 21

Met Leu Lys Ser Ala Leu Leu Leu Gly Ala Ala Ala Val Ser Val Gln

1	5	10	15
Ser Ala Ser Ile Pro Thr Ile Pro Ala Asn Leu Glu Pro Arg Gln Ile			
20	25	30	
Arg Ser Leu Cys Glu Leu Tyr Gly Tyr Trp Ser Gly Asn Gly Tyr Glu			
35	40	45	
Leu Leu Asn Asn Leu Trp Gly Lys Asp Thr Ala Thr Ser Gly Trp Gln			
50	55	60	
Cys Thr Tyr Leu Asp Gly Thr Asn Asn Gly Gly Ile Gln Trp Asn Thr			
65	70	75	80
Ala Trp Glu Trp Gln Gly Ala Pro Asp Asn Val Lys Asn Tyr Pro Tyr			
85	90	95	
Val Gly Lys Gln Ile Gln Arg Gly Arg Lys Ile Ser Asp Ile Asn Ser			
100	105	110	
Met Arg Thr Ser Val Ser Trp Thr Tyr Asp Arg Thr Asp Leu Arg Ala			
115	120	125	
Asn Val Ala Tyr Asp Val Phe Thr Ala Arg Asp Pro Asp His Pro Asn			
130	135	140	
Trp Gly Gly Asp Tyr Glu Leu Met Ile Trp Leu Ala Arg Tyr Gly Gly			
145	150	155	160
Ile Tyr Pro Ile Gly Thr Phe His Ser Gln Val Asn Leu Ala Gly Arg			
165	170	175	
Thr Trp Asp Leu Trp Thr Gly Tyr Asn Gly Asn Met Arg Val Tyr Ser			
180	185	190	
Phe Leu Pro Pro Ser Gly Asp Ile Arg Asp Phe Ser Cys Asp Ile Lys			
195	200	205	
Asp Phe Phe Asn Tyr Leu Glu Arg Asn His Gly Tyr Pro Ala Arg Glu			
210	215	220	
Gln Asn Leu Ile Val Tyr Gln Val Gly Thr Glu Cys Phe Thr Gly Gly			
225	230	235	240
Pro Ala Arg Phe Thr Cys Arg Asp Phe Arg Ala Asp Leu Trp			
245	250		

<210> 22

<211> 247

<212> PRT

<213> Chaetomium brasiliense

<400> 22

Met Lys Leu Thr Leu Val Leu Phe Val Ser Ser Leu Ala Ala Ala Thr	
1	5
Pro Leu Gly Trp Arg Glu Arg Arg Gln Gln Val Ser Leu Cys Gly Gln	
20	25
Ser Ser Ser Trp Ser Gly Asn Gly Tyr Gln Leu Asn Asn Asn Leu Trp	
35	40
Gly Gln Ser Arg Ala Thr Ser Gly Ser Gln Cys Thr Tyr Leu Asp Ser	
50	55
Ser Ser Asn Ser Gly Ile His Trp His Thr Thr Trp Thr Trp Glu Gly	
65	70
Gly Glu Gly Glu Val Lys Ser Tyr Ala Tyr Ser Gly Arg Gln Val Ser	
85	90
Thr Gly Leu Thr Ile Ala Ser Ile Asp Ser Met Gln Thr Ser Val Ser	
100	105
Trp Glu Tyr Asn Thr Thr Asp Ile Gln Ala Asn Val Ala Tyr Asp Ile	
115	120
Phe Thr Ala Glu Asp Pro Asp His Glu His Ser Ser Gly Asp Tyr Glu	
130	135
Val Met Ile Trp Leu Ala Arg Tyr Asn Asn Val Ser Pro Ile Gly Ser	
140	145

145		150		155		160									
Ser	Val	Ala	Thr	Ala	Thr	Val	Gly	Gly	Asp	Thr	Trp	Asp	Leu	Phe	Ala
		165							170					175	
Gly	Ala	Asn	Gly	Asp	Met	Glu	Val	Tyr	Ser	Phe	Val	Ala	Glu	Asn	Thr
		180							185					190	
Met	Asn	Ser	Phe	Ser	Gly	Asp	Val	Lys	Asp	Phe	Phe	Asp	Tyr	Leu	Glu
		195					200					205			
Gln	Asn	Val	Gly	Phe	Pro	Val	Asp	Asp	Gln	Tyr	Leu	Leu	Val	Phe	Glu
	210					215					220				
Leu	Gly	Ser	Glu	Ala	Phe	Thr	Gly	Gly	Pro	Ala	Thr	Leu	Ser	Val	Ser
225					230					235					240
Gln	Phe	Ser	Ala	Asn	Ile	Ala									
				245											

<210> 23

<211> 357

<212> PRT

<213> Bionectria ochroleuca

<400> 23

Met	Lys	Ser	Ile	Ile	Ser	Phe	Phe	Gly	Leu	Ala	Thr	Leu	Val	Ala	Ala
1				5					10					15	
Ala	Pro	Ser	Gln	Asn	Pro	Thr	Arg	Thr	Gln	Pro	Leu	Glu	Lys	Arg	Ala
			20					25					30		
Thr	Thr	Leu	Cys	Gly	Gln	Trp	Asp	Ser	Val	Glu	Thr	Gly	Gly	Tyr	Thr
		35					40					45			
Ile	Tyr	Asn	Asn	Leu	Trp	Gly	Gln	Asp	Asn	Gly	Ser	Gly	Ser	Gln	Cys
			50			55					60				
Leu	Thr	Val	Glu	Gly	Val	Thr	Asp	Gly	Leu	Ala	Ala	Trp	Ser	Ser	Thr
65					70					75					80
Trp	Ser	Trp	Ser	Gly	Gly	Ser	Ser	Ser	Val	Lys	Ser	Tyr	Ser	Asn	Ala
			85						90					95	
Val	Leu	Ser	Ala	Glu	Ala	Ala	Arg	Ile	Ser	Ala	Ile	Ser	Ser	Ile	Pro
			100					105					110		
Ser	Lys	Trp	Glu	Trp	Ser	Tyr	Thr	Gly	Thr	Asp	Ile	Val	Ala	Asn	Val
		115					120					125			
Ala	Tyr	Asp	Leu	Phe	Ser	Asn	Thr	Asp	Cys	Gly	Asp	Thr	Pro	Glu	Tyr
	130					135					140				
Glu	Ile	Met	Ile	Trp	Leu	Ser	Ala	Leu	Gly	Gly	Ala	Gly	Pro	Ile	Ser
145					150					155					160
Ser	Thr	Gly	Ser	Ser	Ile	Ala	Thr	Val	Thr	Ile	Ala	Gly	Ala	Ser	Trp
			165						170					175	
Asn	Leu	Trp	Gln	Gly	Gln	Asn	Asn	Gln	Met	Thr	Val	Phe	Ser	Phe	Val
			180					185					190		
Ala	Glu	Ser	Asp	Gln	Lys	Ser	Phe	Ser	Gly	Asp	Leu	Asn	Asp	Phe	Ile
		195					200					205			
Gln	Tyr	Leu	Val	Asp	Ser	Gln	Gly	Tyr	Ser	Gly	Ser	Gln	Cys	Leu	Tyr
	210					215					220				
Ser	Ile	Gly	Ala	Gly	Thr	Glu	Pro	Phe	Thr	Gly	Thr	Asp	Ala	Glu	Phe
225					230					235					240
Ile	Thr	Thr	Gly	Tyr	Ser	Val	Ser	Val	Ser	Ala	Gly	Asp	Ser	Gly	Ser
			245						250					255	
Asp	Glu	Thr	Thr	Ser	Ser	Gln	Ala	Gln	Ser	Ser	Thr	Val	Glu	Thr	
			260				265						270		
Ser	Thr	Ala	Thr	Gln	Pro	Gln	Ser	Ser	Ser	Thr	Val	Val	Pro	Thr	Val
		275					280					285			

Thr Leu Ser Gln Pro Ser Asn Glu Ser Thr Thr Thr Pro Val Gln Ser
 290 295 300
 Gln Pro Ser Ser Val Glu Thr Thr Pro Thr Ala Gln Pro Gln Ser Ser
 305 310 315 320
 Ser Val Gln Thr Thr Thr Thr Ala Gln Ala Gln Pro Thr Pro Glu Arg
 325 330 335
 Ala Ala Pro Asp Ala Gly Ser Ala Glu Leu Leu Ser Ser Ala Thr Met
 340 345 350
 His Leu Asp Arg Arg
 355

<210> 24
 <211> 247
 <212> PRT
 <213> *Emericella desertorum*

<400> 24
 Met Lys Leu Leu Ala Leu Ser Leu Val Ser Leu Ala Ser Ala Ala Ser
 1 5 10 15
 Ala Ala Ser Ile Leu Ser Asn Thr Phe Thr Arg Arg Ser Asp Phe Cys
 20 25 30
 Gly Gln Trp Asp Thr Ala Thr Val Gly Asn Phe Ile Val Tyr Asn Asn
 35 40 45
 Leu Trp Gly Gln Asp Asn Ala Asp Ser Gly Ser Gln Cys Thr Gly Val
 50 55 60
 Asp Ser Ala Asn Gly Asn Ser Ile Ser Trp His Thr Thr Trp Ser Trp
 65 70 75 80
 Ser Gly Gly Ser Ser Ser Val Lys Ser Tyr Ala Asn Ala Ala Tyr Gln
 85 90 95
 Phe Thr Ser Thr Lys Leu Asn Ser Leu Ser Ser Ile Pro Thr Ser Trp
 100 105 110
 Lys Trp Gln Tyr Ser Thr Thr Asp Ile Val Ala Asn Val Ala Tyr Asp
 115 120 125
 Leu Phe Thr Ser Ser Ser Ala Gly Gly Asp Ser Glu Tyr Glu Ile Met
 130 135 140
 Ile Trp Leu Ala Ala Leu Gly Gly Ala Gly Pro Ile Ser Ser Thr Gly
 145 150 155 160
 Ser Ser Ile Ala Thr Val Thr Leu Gly Gly Val Thr Trp Ser Leu Tyr
 165 170 175
 Ser Gly Pro Asn Gly Ser Met Gln Val Tyr Ser Phe Val Ala Ser Ser
 180 185 190
 Thr Thr Glu Ser Phe Ser Ala Asp Leu Met Asp Phe Ile Asn Tyr Leu
 195 200 205
 Ala Glu Asn Gln Gly Leu Ser Ser Ser Gln Tyr Leu Thr His Val Gln
 210 215 220
 Ala Gly Thr Glu Pro Phe Thr Gly Thr Asp Ala Thr Leu Thr Val Ser
 225 230 235 240
 Ser Tyr Ser Val Ser Val Ser
 245

<210> 25
 <211> 244
 <212> PRT
 <213> *Fusarium solani*

<400> 25
 Met Lys Ser Ala Ile Val Ala Ala Leu Ala Gly Leu Ala Ala Ala Ser

1				5					10					15			
Pro	Thr	Arg	Leu	Ile	Pro	Arg	Gly	Gln	Phe	Cys	Gly	Gln	Trp	Asp	Ser		
			20					25					30				
Glu	Thr	Ala	Gly	Ala	Tyr	Thr	Ile	Tyr	Asn	Asn	Leu	Trp	Gly	Lys	Asp		
		35					40					45					
Asn	Ala	Glu	Ser	Gly	Glu	Gln	Cys	Thr	Thr	Asn	Ser	Gly	Glu	Gln	Ser		
	50					55					60						
Asp	Gly	Ser	Ile	Ala	Trp	Ser	Val	Glu	Trp	Ser	Trp	Thr	Gly	Gly	Gln		
65					70					75					80		
Gly	Gln	Val	Lys	Ser	Tyr	Pro	Asn	Ala	Val	Val	Glu	Ile	Glu	Lys	Lys		
			85					90					95				
Thr	Leu	Gly	Glu	Val	Ser	Ser	Ile	Pro	Ser	Ala	Trp	Asp	Trp	Thr	Tyr		
		100						105					110				
Thr	Gly	Asn	Gly	Ile	Ile	Ala	Asn	Val	Ala	Tyr	Asp	Leu	Phe	Thr	Ser		
		115					120					125					
Ser	Thr	Glu	Ser	Gly	Asp	Ala	Glu	Tyr	Glu	Phe	Met	Ile	Trp	Leu	Ser		
	130					135					140						
Ala	Leu	Gly	Gly	Ala	Gly	Pro	Ile	Ser	Asn	Asp	Gly	Ser	Pro	Val	Ala		
145					150					155					160		
Thr	Val	Glu	Leu	Ala	Gly	Thr	Ser	Trp	Lys	Leu	Tyr	Gln	Gly	Lys	Asn		
				165					170					175			
Asn	Gln	Met	Thr	Val	Phe	Ser	Phe	Val	Ala	Glu	Ser	Asp	Val	Asn	Asn		
			180					185					190				
Phe	Cys	Gly	Asp	Leu	Ala	Asp	Phe	Thr	Asp	Tyr	Leu	Val	Asp	Asn	His		
		195					200					205					
Gly	Val	Ser	Ser	Ser	Gln	Ile	Leu	Gln	Ser	Val	Gly	Ala	Gly	Thr	Glu		
	210					215					220						
Pro	Phe	Glu	Gly	Thr	Asn	Ala	Val	Phe	Thr	Thr	Asn	Asn	Tyr	His	Ala		
225					230					235					240		
Asp	Val	Glu	Tyr														

<210> 26

<211> 250

<212> PRT

<213> Fusarium solani

<400> 26

Met	Lys	Phe	Phe	Gly	Val	Val	Ser	Ala	Phe	Leu	Ala	Ala	Thr	Ala	Val		
1				5					10					15			
Ala	Thr	Pro	Thr	Thr	Pro	Thr	Glu	Thr	Ile	Glu	Lys	Arg	Asp	Thr	Thr		
			20					25					30				
Trp	Cys	Asp	Ala	Phe	Gly	Ser	Leu	Ala	Thr	Ser	Gly	Tyr	Thr	Val	Tyr		
		35					40					45					
His	Asn	Asn	Trp	Gly	Lys	Gly	Asp	Ala	Thr	Ser	Gly	Ser	Gln	Cys	Thr		
	50					55					60						
Thr	Phe	Thr	Ser	Val	Ser	Asn	Asn	Asn	Phe	Val	Trp	Ser	Thr	Ser	Trp		
65					70					75					80		
Thr	Trp	Ala	Gly	Gly	Ala	Gly	Lys	Val	Lys	Ser	Tyr	Ser	Asn	Val	Ala		
			85						90					95			
Leu	Glu	Lys	Ile	Asn	Lys	Lys	Ile	Ser	Asp	Ile	Lys	Ser	Val	Ser	Thr		
			100					105					110				
Arg	Trp	Ile	Trp	Arg	Tyr	Thr	Gly	Thr	Lys	Met	Ile	Ala	Asn	Val	Ser		
		115					120					125					
Tyr	Asp	Leu	Trp	Phe	Ala	Pro	Thr	Ala	Ser	Ser	Asn	Asn	Ala	Tyr	Glu		
	130					135					140						
Ile	Met	Ile	Trp	Val	Gly	Ala	Tyr	Gly	Gly	Ala	Leu	Pro	Ile	Ser	Thr		
145					150					155					160		

Pro	Gly	Lys	Gly	Val	Ile	Asp	Arg	Pro	Thr	Leu	Ala	Gly	Ile	Pro	Trp
				165					170					175	
Asp	Val	Tyr	Lys	Gly	Pro	Asn	Gly	Asp	Val	Thr	Val	Ile	Ser	Phe	Val
			180					185						190	
Ala	Ser	Ser	Asn	Gln	Gly	Asn	Phe	Gln	Ala	Asp	Leu	Lys	Glu	Phe	Leu
		195					200					205			
Asn	Tyr	Leu	Thr	Ser	Lys	Gln	Gly	Leu	Pro	Ser	Asn	Tyr	Val	Ala	Thr
	210					215					220				
Ser	Phe	Gln	Ala	Gly	Thr	Glu	Pro	Phe	Glu	Gly	Thr	Asn	Ala	Val	Leu
225					230					235					240
Lys	Thr	Ser	Ala	Tyr	Thr	Ile	Ser	Val	Asn						
				245					250						

<210> 27

<211> 371

<212> PRT

<213> Streptomyces sp. 11AG8

<400> 27

Met	Arg	Ser	His	Pro	Arg	Ser	Ala	Thr	Met	Thr	Val	Leu	Val	Val	Leu
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Ala	Ser	Leu	Gly	Ala	Leu	Leu	Thr	Ala	Ala	Ala	Pro	Ala	Gln	Ala	Asn
			20					25					30		
Gln	Gln	Ile	Cys	Asp	Arg	Tyr	Gly	Thr	Thr	Thr	Ile	Gln	Asp	Arg	Tyr
		35					40					45			
Val	Val	Gln	Asn	Asn	Arg	Trp	Gly	Thr	Ser	Ala	Thr	Gln	Cys	Ile	Asn
	50					55					60				
Val	Thr	Gly	Asn	Gly	Phe	Glu	Ile	Thr	Gln	Ala	Asp	Gly	Ser	Val	Pro
65					70				75						80
Thr	Asn	Gly	Ala	Pro	Lys	Ser	Tyr	Pro	Ser	Val	Tyr	Asp	Gly	Cys	His
				85					90					95	
Tyr	Gly	Asn	Cys	Ala	Pro	Arg	Thr	Thr	Leu	Pro	Met	Arg	Ile	Ser	Ser
			100					105					110		
Ile	Gly	Ser	Ala	Pro	Ser	Ser	Val	Ser	Tyr	Arg	Tyr	Thr	Gly	Asn	Gly
		115					120					125			
Val	Tyr	Asn	Ala	Ala	Tyr	Asp	Ile	Trp	Leu	Asp	Pro	Thr	Pro	Arg	Thr
	130					135					140				
Asn	Gly	Val	Asn	Arg	Thr	Glu	Ile	Met	Ile	Trp	Phe	Asn	Arg	Val	Gly
145					150					155					160
Pro	Val	Gln	Pro	Ile	Gly	Ser	Pro	Val	Gly	Thr	Ala	His	Val	Gly	Gly
				165					170					175	
Arg	Ser	Trp	Glu	Val	Trp	Thr	Gly	Ser	Asn	Gly	Ser	Asn	Asp	Val	Ile
			180				185						190		
Ser	Phe	Leu	Ala	Pro	Ser	Ala	Ile	Ser	Ser	Trp	Ser	Phe	Asp	Val	Lys
	195						200					205			
Asp	Phe	Val	Asp	Gln	Ala	Val	Ser	His	Gly	Leu	Ala	Thr	Pro	Asp	Trp
	210					215					220				
Tyr	Leu	Thr	Ser	Ile	Gln	Ala	Gly	Phe	Glu	Pro	Trp	Glu	Gly	Gly	Thr
225					230					235					240
Gly	Leu	Ala	Val	Asn	Ser	Phe	Ser	Ser	Ala	Val	Asn	Ala	Gly	Gly	Gly
				245					250					255	
Asn	Gly	Gly	Thr	Pro	Gly	Thr	Pro	Ala	Ala	Cys	Gln	Val	Ser	Tyr	Ser
			260					265					270		
Thr	His	Thr	Trp	Pro	Gly	Gly	Phe	Thr	Val	Asp	Thr	Thr	Ile	Thr	Asn
	275						280					285			
Thr	Gly	Ser	Thr	Pro	Val	Asp	Gly	Trp	Glu	Leu	Asp	Phe	Thr	Leu	Pro
	290					295					300				

Ala Gly His Thr Val Thr Ser Val Trp Asn Ala Leu Ile Ser Pro Ala
 305 310 315 320
 Ser Gly Ala Val Thr Ala Arg Ser Thr Gly Ser Asn Gly Arg Ile Ala
 325 330 335
 Ala Asn Gly Gly Thr Gln Ser Phe Gly Phe Gln Gly Thr Ser Ser Gly
 340 345 350
 Ala Gly Phe Thr Ala Pro Ala Gly Ala Arg Leu Asn Gly Thr Ser Cys
 355 360 365
 Thr Val Arg
 370

<210> 28
 <211> 221
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> consensus sequence

<220>
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 <222> (1)...(221)
 <223> Xaa = Any Amino Acid

<400> 28
 Cys Xaa Gln Tyr Xaa Xaa Xaa Xaa Xaa Xaa Gly Tyr Xaa Xaa Xaa Asn
 1 5 10 15
 Asn Xaa Trp Gly Xaa Xaa Xaa Xaa Xaa Ser Gly Xaa Gln Cys Thr Xaa
 20 25 30
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Trp Xaa Xaa Xaa Trp
 35 40 45
 Xaa Trp Ser Gly Gly Xaa Xaa Xaa Val Lys Ser Tyr Xaa Xaa Xaa Xaa
 50 55 60
 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ser Xaa Ile Xaa Ser Xaa
 65 70 75 80
 Xaa Xaa Xaa Xaa Xaa Trp Xaa Tyr Xaa Gly Xaa Xaa Xaa Xaa Ala Asn
 85 90 95
 Val Ala Tyr Asp Leu Phe Thr Xaa Xaa Xaa Pro Xaa His Xaa Xaa Xaa
 100 105 110
 Xaa Gly Xaa Tyr Glu Xaa Met Ile Trp Leu Xaa Xaa Xaa Gly Gly Xaa
 115 120 125
 Xaa Pro Ile Gly Ser Xaa Xaa Xaa Xaa Val Xaa Xaa Xaa Xaa Xaa Xaa
 130 135 140
 Gly Xaa Xaa Trp Xaa Leu Xaa Xaa Gly Xaa Asn Gly Xaa Met Xaa Val
 145 150 155 160
 Xaa Ser Phe Val Ala Xaa Ser Ser Ser Ser Ser Ser Phe Xaa Gly Asp
 165 170 175
 Xaa Xaa Xaa Phe Xaa Xaa Tyr Leu Xaa Xaa Xaa Xaa Gly Xaa Pro Xaa
 180 185 190
 Xaa Xaa Gln Tyr Leu Xaa Xaa Xaa Gln Xaa Gly Thr Glu Pro Phe Thr
 195 200 205
 Gly Xaa Xaa Ala Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Ala
 210 215 220

<210> 29
 <211> 25
 <212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 29

gaacgatggc aagggcggcg tgacg

25

<210> 30

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 30

cttctcgggc tgctacaacc caaacgg

27

<210> 31

<211> 26

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 31

acatcgtcga gtgttttggc acctac

26

<210> 32

<211> 29

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 32

catcgtcag aactggggca cctacaacc

29

<210> 33

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 33

ggcacctacc gaccgtccac g

21

<210> 34

<211> 25

<212> DNA

<213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

 <400> 34
 caagctgggc gagcacacct ccgac 25

 <210> 35
 <211> 21
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <400> 35
 cgccgcaact gtcgctcgag c 21

 <210> 36
 <211> 29
 <212> DNA
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 <220>
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 <400> 36
 gtggagggtt accaaagctc tggctctgc 29

 <210> 37
 <211> 27
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 <400> 37
 tctggctctg cttgcatcac cgtcagc 27

 <210> 38
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 <212> DNA
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 <223> synthetic oligonucleotide

 <400> 38
 gagaagcgcc agtgcattca gcccggc 27

 <210> 39
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 <212> DNA
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<400> 39

gtgacgtact gcaatgggtcc cggcggg

27

<210> 40

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 40

ggcaccaaga acaggggtcat caacttctcg ggc

33

<210> 41

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 41

tccatcaccg tcagcgatta aagggggctc ttc

33

<210> 42

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 42

cccagacgat tcagtgcggc acgggctaca ac

32

<210> 43

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 43

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32

<210> 44

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> synthetic oligonucleotide

<400> 44
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<210> 45
 <211> 35
 <212> DNA
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<220>
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<400> 45
 caacggctac ttctactgct actggaacga tggcc 35

<210> 46
 <211> 34
 <212> DNA
 <213> Artificial Sequence

<220>
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<400> 46
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<210> 47
 <211> 31
 <212> DNA
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<220>
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<400> 47
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<210> 48
 <211> 27
 <212> DNA
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<220>
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<400> 48
 ggcgccacca agtgcggcga ggtcacc 27

<210> 49
 <211> 28
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> synthetic oligonucleotide

<400> 49

gcgtgggctc agtgcggcct gacgctcg

28

<210> 50

<211> 752

<212> DNA

<213> *Trichoderma reesei*

<400> 50

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cccacaggcc	tgcagcctga	gagcagtgtc	aacgtcacag	agcgtggcat	gtacgacttt	120
gttcttggag	ctcacaatga	tcctgcgcgt	cgtgctagca	tcaactacga	ccaaaactac	180
caaactggcg	gacaagtcag	ctattcgcc	tccaacactg	gcttctcagt	gaactggaac	240
actcaagatg	actttgttgt	gggcgttggt	tggacgactg	gatcttctgc	gtaggaggac	300
tcctcatcat	tctgcacttt	gaaagcatct	tctgaccaa	agcttctctt	agtcctcatc	360
actttggcgg	ctcttttagt	gtcaacagcg	gaactggcct	gctttccgtc	tatggctgga	420
gcaccaaccc	actggttgag	tactacatca	tggaggacaa	ccacaactac	ccagcacagg	480
gtaccgtcaa	gggaaccgtc	accagcgacg	gagccactta	caccatctgg	gagaataccc	540
gtgtcaacga	gccttccatc	cagggcacag	cgaccttcaa	ccagtacatt	tccgtgcgga	600
actgcgccag	gaccagcgga	actgttactg	tgcagaacca	cttcaatgct	tgggcctcgc	660
ttggcctgca	ccttgggcag	atgaactacc	aggttgctgc	tgtcgaaggc	tggggtggtg	720
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<210> 51

<211> 248

<212> PRT

<213> *Trichoderma reesei*

<400> 51

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Thr	Leu	Ala	Met	Pro	Thr	Gly	Leu	Glu	Pro	Glu	Ser	Ser	Val	Asn
		20					25					30		Val
Thr	Glu	Arg	Gly	Met	Tyr	Asp	Phe	Val	Leu	Gly	Ala	His	Asn	Asp
		35				40					45			His
Arg	Arg	Arg	Ala	Ser	Ile	Asn	Tyr	Asp	Gln	Asn	Tyr	Gln	Thr	Gly
		50				55					60			Gly
Gln	Val	Ser	Tyr	Ser	Pro	Ser	Asn	Thr	Gly	Phe	Ser	Val	Asn	Trp
					70					75				Asn
Thr	Gln	Asp	Asp	Phe	Val	Val	Gly	Val	Gly	Trp	Thr	Thr	Gly	Ser
				85					90					Ser
Ala	Glu	Asp	Ser	Ser	Ser	Phe	Cys	Thr	Leu	Lys	Ala	Ser	Ser	Asp
			100					105					110	Gln
Lys	Leu	Leu	Leu	Val	Pro	Ser	Thr	Leu	Ala	Ala	Leu	Leu	Val	Ser
		115					120					125		Thr
Ala	Glu	Leu	Ala	Cys	Phe	Pro	Ser	Met	Ala	Gly	Ala	Pro	Thr	His
		130				135					140			Trp
Leu	Ser	Thr	Thr	Ser	Trp	Arg	Thr	Thr	Thr	Thr	Thr	Gln	His	Arg
					150					155				Val
Pro	Ser	Arg	Glu	Pro	Ser	Pro	Ala	Thr	Glu	Pro	Leu	Thr	Pro	Ser
			165						170					Gly
Arg	Ile	Pro	Val	Ser	Thr	Ser	Leu	Pro	Ser	Arg	Ala	Gln	Arg	Pro
			180					185					190	Ser
Thr	Ser	Thr	Phe	Pro	Cys	Gly	Thr	Arg	Pro	Gly	Pro	Ala	Glu	Leu
		195					200					205		Leu
Leu	Cys	Arg	Thr	Thr	Ser	Met	Leu	Gly	Pro	Arg	Leu	Ala	Cys	Thr
		210				215					220			Leu
Gly	Arg	Thr	Thr	Arg	Leu	Ser	Leu	Ser	Lys	Ala	Gly	Val	Val	Val

225
Leu Pro His Arg Val Ser Ala Thr
245

230

235

240